

**Amendments to the Claims:**

Please amend claims 2-7, 10 and 12-13, add new claims 56-60, and cancel claims 1 and 8-9.  
This listing of claims will replace all prior versions, and listings of claims in the application:

**Listing of Claims:**

- 1                    1.        (canceled).
- 1                    2.        (currently amended) The method of claim ~~1~~ 10 wherein the presence or  
2        absence of a nick in a DNA molecule is measured by determining the change in electrophoretic  
3        mobility of nicked DNA on an electrophoretic gel.
- 1                    3.        (currently amended) The method of claim ~~1~~ 10 wherein the presence or  
2        absence of a nick in a DNA molecule is determined by a SI nuclease assay.
- 1                    4.        (currently amended) The method of claim ~~1~~ 10 wherein the presence or  
2        absence of a nick in a DNA molecule is determined by a primer extension reaction.
- 1                    5.        (currently amended) The method of claim ~~1~~ 10 wherein the presence or  
2        absence of a nick in a DNA molecule is determined by a polymerase chain reaction amplification  
3        reaction.
- 1                    6.        (currently amended) The method of claim ~~1~~ 10 wherein the presence or  
2        absence of a nick in a DNA molecule is determined by a DNA sequencing assay.
- 1                    7.        (currently amended) The method of claim ~~1~~ 10 wherein the presence or  
2        absence of a nick in a DNA molecule is determined by a protein binding assay.
- 1                    8.        (canceled).
- 1                    9.        (canceled).

1                   10.   (currently amended) A method of detecting eukaryotic nicking  
2 transcription factor activity comprising the steps of:  
3                   a)     providing a DNA template comprising at least one binding region for a  
4 transcription factor;  
5                   b)     contacting the DNA template with at least one eukaryotic nicking  
6 transcription factor; and  
7                   c)     detecting the presence or absence of a nick in the DNA template at or near  
8 the binding region of the eukaryotic nicking transcription factor, wherein the presence of a nick  
9 in the DNA template indicates nicking transcription factor activity.

1                   11.   (original)     The method of claim 10, wherein the transcription factor is  
2 in a nuclear cell extract.

1                   12.   (currently amended) The method of claim 10, further comprising the  
2 steps of:  
3                   a)     isolating the DNA template; and  
4                   b)     ~~wherein the DNA template is inserted~~ inserting the DNA template into a  
5 viral or plasmid vector and ~~introduced~~ introducing the template into a cell.

1                   13.   (currently amended) The method of claim 10, ~~wherein the DNA template~~  
2 ~~is fixed to a matrix~~, further comprising the step of fixing the DNA template to a matrix.

1                   14.   (original)     The method of claim 13, wherein the matrix is a biological  
2 chip.

1                   15-55. (canceled).

1                   56.   (new)     The method of claim 10, wherein the eukaryotic nicking  
2 transcription factor comprises a site specific DNA binding transcription factor.

1                    57.    (new) The method of claim 56, wherein the eukaryotic nicking  
2 transcription factor comprises an enhancer binding protein.

1                    58.    (new) The method of claim 56, wherein the eukaryotic nicking  
2 transcription factor comprises a general transcription factor.

1                    59.    (new) The method of claim 10, wherein the eukaryotic nicking  
2 transcription factor regulates the transcription of structural RNA.

1                    60.    (new) The method of claim 10, wherein the eukaryotic nicking  
2 transcription factor regulates the transcription of protein encoding RNA.

1                    61.    (new) The method of claim 10, wherein the eukaryotic nicking  
2 transcription factor is selected from the group consisting of CREB, TFIIC and c-jun.